BIOTECHNOLOGY SYSTEMS BRANCH

MAY 1 0 2002 MAY 1 0 2002 TER 160069

RAW SEQUENCE LISTING ERROR REPORT

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Application Serial Number:	09/425,804B
Source:	1600
Date Processed by STIC:	5/1/02

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

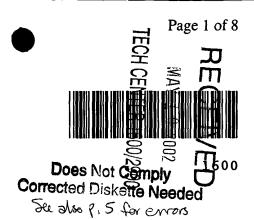
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- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
- Hand Carry directly to:
 U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
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Revised 01/29/2002



RAW SEQUENCE LISTING

22 <170> SOFTWARE: PatentIn Ver. 2.1

DATE: 05/01/2002 TIME: 12:34:25

PATENT APPLICATION: US/09/425,804B

Input Set : A:\09425804_Ref No 122144-1001 SEQ ID LISTING.txt

Output Set: N:\CRF3\05012002\I425804B.raw

4 <110> APPLICANT: Gorenstein, David G. Luxon, Bruce A. Herzog, Norbert 6 Aronson, Judy 9 <120> TITLE OF INVENTION: Thio-Modified Aptamer Synthetic Methods and Compositions 10 12 <130> FILE REFERENCE: 122144-001001 14 <140> CURRENT APPLICATION NUMBER: 09/425,804B 15 <141> CURRENT FILING DATE: 1999-10-25 17 <150> PRIOR APPLICATION NUMBER: 60/105,600 18 <151> PRIOR FILING DATE: 1998-10-26 20 <160> NUMBER OF SEQ ID NOS: 50

ERRORED SEQUENCES

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414 <400> SEQUENCE: 33

DATE: 05/01/2002

PATENT APPLICATION: US/09/425,804B TIME: 12:34:25 Input Set : A:\09425804_Ref No 122144-1001 SEQ ID LISTING.txt Output Set: N:\CRF3\05012002\I425804B.raw (14) - line count is 15 E--> 415 gtgggaatctc ctgg 435 <210> SEQ ID NO: 35 436 <211> LENGTH: 14 437 <212> TYPE: DNA 438 <213> ORGANISM: Artificial Sequence 440 <220> FEATURE: 441 <221> NAME/KEY: modified_base W--> 442 (212) TYPE: (2) change to 22227 - Only needed once E--> 442 (212) TYPE: (2) delete W--> 443 <213> ORGANISM: thymidine 3'-O-phosphorothioate 445 <220> FEATURE: 446 <221> NAME/KEY: modified_base 447 <222> LOCATION: (7) 448 <223> OTHER INFORMATION: thymidine 3'-O-phosphorothioate 450 <220> FEATURE: 451 <221> NAME/KEY: modified_base 452 <222> LOCATION: (9) 453 <223> OTHER INFORMATION: thymidine 3'-O-phosphorothioate 455 <220> FEATURE: 456 <221> NAME/KEY: modified_base 457 <222> LOCATION: (12) 458 <223> OTHER INFORMATION: thymidine 3'-O-phosphorothioate 460 <400> SEQUENCE: 35 461 gtggaatctc ctgg 14 463 <210> SEQ ID NO: 36 464 <211> LENGTH: 14 465 <212> TYPE: DNA 466 <213> ORGANISM: Artificial Sequence 468 <220> FEATURE: 469 <221> NAME/KEY: modified_base - Some error as tobac W--> 470 <212> TYPE: Compliment (2) E--> 470 <212> TYPE: Compliment (2) W--> 471 (213>)ORGANISM: phosphorothioate \rightarrow charge to $\angle 2237$ 473 <220> FEATURE: 474 <221> NAME/KEY: modified_base W--> 475 <212> TYPE: Compliment (12) E--> 475 <212> TYPE+ Compliment (12) W--> 476 (213) ORGANISM: phosphorothioate 478 <220> FEATURE: 479 <223> OTHER INFORMATION: Description of Artificial Sequence: aptamer 481 <400> SEQUENCE: 36 482 ccaggagatt ccac 14 484 <210> SEQ ID NO: 37 485 <211> LENGTH: 14 486 <212> TYPE: DNA

RAW SEQUENCE LISTING

489 <220> FEATURE:

W--> $491 \langle 212 \rangle$ TYPE: Compliment (5)

42227

487 <213> ORGANISM: Artificial Sequence

490 <221> NAME/KEY: modified_base

RAW SEQUENCE LISTING

DATE: 05/01/2002

PATENT APPLICATION: US/09/425,804B

TIME: 12:34:25

Input Set : A:\09425804_Ref No 122144-1001 SEQ ID LISTING.txt

Output Set: N:\CRF3\05012002\I425804B.raw

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E--> 491 (212) TYPE: Compliment (5) delete
W--> 492 (213) ORGANISM: phosphorothioate
                                               -3 22235
     494 <220> FEATURE:
495 <221> NAME/KEY: modified_base
W--> 496 (212) TYPE: Compliment (7) | 2227
E--> 496 (212) TYPE: Compliment (7) delete
W--> 497 (213) ORGANISM: phosphorothicate > <223>
     499 <220> FEATURE:
     500 <223> OTHER INFORMATION: Description of Artificial Sequence: aptamer
     502 <400> SEQUENCE: 37
     503 ccaggagatt ccac
                                  14
     562 <210> SEQ ID NO: 40
     563 <211> LENGTH: 14
     564 <212> TYPE: DNA
     565 <213> ORGANISM: Artificial Sequence
     567 <220> FEATURE:
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     569 <222> LOCATION: (9)..(10)
     570 <223> OTHER INFORMATION: phosphorothioate
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     578 <221> NAME/KEY: modified_base
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     580 <223> OTHER INFORMATION: phosphorothioate
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     583 <223> OTHER INFORMATION: Description of Artificial Sequence: aptamer
E--> 585 <400> SEQUENCE: 38
                              Dupped sequence, see P.S
     586 ccaggagatt ccac
E--> 588 <210> SEQ ID NO: 40
     589 <211> LENGTH: 14
     590 <212> TYPE: DNA
     591 <213> ORGANISM: Artificial Sequence
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     594 <221> NAME/KEY: modified_base
     595 <222> LOCATION: (9)..(10)
     596 <223> OTHER INFORMATION: phosphorothioate
     598 <220> FEATURE:
     599 <221> NAME/KEY: modified_base
     600 <222> LOCATION: Complement (2)
     601 <223> OTHER INFORMATION: phosphorothioate
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     604 <221> NAME/KEY: modified_base
     605 <222> LOCATION: Complement (5)
     606 <223> OTHER INFORMATION: phosphorothioate
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608 <220> FEATURE:

609 <221> NAME/KEY: modified_base

RAW SEQUENCE LISTING

DATE: 05/01/2002 TIME: 12:34:25

PATENT APPLICATION: US/09/425,804B

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Input Set: A:\09425804_Ref No 122144-1001 SEQ ID LISTING.txt

Output Set: N:\CRF3\05012002\1425804B.raw

610 <222> LOCATION: Complement (7) 611 <223> OTHER INFORMATION: phosphorothioate 613 <220> FEATURE: 614 <221> NAME/KEY: modified_base 615 <222> LOCATION: Complement (12) 616 <223> OTHER INFORMATION: phosphorothioate 618 <220> FEATURE: 619 <223> OTHER INFORMATION: Description of Artificial Sequence: aptamer E--> 621 <400> SEQUENCE: (41) - NO. differs from Seq. ID No. 622 ccaggagatt ccac E--> 624 <210> SEQ ID NO: (42) — Should be 41 622 ccaggagatt ccac 14 635 <210> SEQ ID NO: 43 636 <211> LENGTH: 43 637 <212> TYPE: DNA 638 <213> ORGANISM: Artificial Sequence 640 <220> FEATURE: 641 <221> NAME/KEY: modified_base W--> 642 (212) TYPE: (9) E--> 642 (212> TYPE: (9) delett W--> 643 (213) ORGANISM: phosphorodithicate > 22237 645 <220> FEATURE: 646 <221> NAME/KEY: modified_base W--> 647 (212) TYPE: (20) --->£2227 E--> 647 (212> TYPE: (20) delete W--> 648 (213) ORGANISM: phosphorodithioate 650 <220> FEATURE: うくひろう 651 <221> NAME/KEY: modified_base W--> 652 (212) TYPE: (9) -> K2227 E--> 652 <212> TYPE+ (9) W--> 653 (213) ORGANISM: phosphate ester 655 <220> FEATURE:

-->∠222>

656 <221> NAME/KEY: modified_base

E--> 661 ccaggagatt ccacttttgt ggaatctcct gga

W--> 658 (213) ORGANISM: phosphate ester

W--> 657 (212) TYPE: (20)

E--> 657- (212> TYPE+ (20) delek

w--> 660 ₹223> OTHER INFORMATION: 660 <400> SEQUENCE: 43

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/425,804B

DATE: 05/01/2002 TIME: 12:34:26

Input Set : A:\09425804_Ref No 122144-1001 SEQ ID LISTING.txt

Output Set: N:\CRF3\05012002\I425804B.raw

Skipped Sequences(NEW RULES):

Sequence(s)_missing. If intentional, please use the following format for each skipped sequence.
<210> sequence id number

<400> sequence id number

Seq#:41

Use of <220> Feature(NEW RULES):

Sequence(s)_are missing the <220> Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register," 6/01/98, Vol. 63, No. 104,pp.29631-32) (Sec.1.823 of new Rules)

Seq#:2,14,25,43

VERIFICATION SUMMARY DATE: 05/01/2002 PATENT APPLICATION: US/09/425,804B TIME: 12:34:26

Input Set: A:\09425804_Ref No 122144-1001 SEQ ID LISTING.txt
Output Set: N:\CRF3\05012002\1425804B.raw

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L:38 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:38 M:254 E: No. of Bases conflict, LENGTH:Input:60 Counted:59 SEQ:1
M:254 Repeated in SeqNo=1
L:39 M:252 E: No. of Seq. differs, <211> LENGTH:Input:66 Found:65 SEQ:1
L:46 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
L:48 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:159 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:176 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14
L:178 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:178 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:258 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:298 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:298 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:317 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0
L:357 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0
L:415 M:254 E: No. of Bases conflict, LENGTH:Input:14 Counted:15 SEQ:33
L:415 M:252 E: No. of Seq. differs, <211> LENGTH:Input:14 Found:15 SEQ:33
L:442 M:280 W: Numeric Identifier already exists, Type not replaced.
L:442 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:443 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:470 M:280 W: Numeric Identifier already exists, Type not replaced.
L:470 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:471 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:475 M:280 W: Numeric Identifier already exists, Type not replaced.
L:475 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:476 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:491 M:280 W: Numeric Identifier already exists, Type not replaced.
L:491 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:492 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:496 M:280 W: Numeric Identifier already exists, Type not replaced.
L:496 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:497 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:585 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:40 differs:38
L:588 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQ ID NO:40
L:621 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:40 differs:41
L:624 M:214 E: (33) Seq.# missing, SEQ ID NO:41
L:642 M:280 W: Numeric Identifier already exists, Type not replaced.
L:642 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:643 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:647 M:280 W: Numeric Identifier already exists, Type not replaced.
L:647 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:648 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:652 M:280 W: Numeric Identifier already exists, Type not replaced.
L:652 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:653 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:657 M:280 W: Numeric Identifier already exists, Type not replaced.
L:657 M:310 E: (3) Wrong or Missing Sequence Type, TYPE:
L:658 M:280 W: Numeric Identifier already exists, Organism not replaced.
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VERIFICATION SUMMARY

DATE: 05/01/2002

PATENT APPLICATION: US/09/425,804B

TIME: 12:34:26

Input Set : A:\09425804_Ref No 122144-1001 SEQ ID LISTING.txt

Output Set: N:\CRF3\05012002\I425804B.raw

 $L:660\ M:258\ W:$ Mandatory Feature missing, <223> OTHER INFORMATION:

L:661 M:254 E: No. of Bases conflict, LENGTH:Input:43 Counted:33 SEQ:43 L:661 M:252 E: No. of Seq. differs, <211> LENGTH:Input:43 Found:33 SEQ:43